



## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification <sup>6</sup> : <b>C12N 15/11, C07H 21/04, A61K 31/70</b>		<b>A2</b>	(11) International Publication Number: <b>WO 98/33904</b>
			(43) International Publication Date: 6 August 1998 (06.08.98)
(21) International Application Number: <b>PCT/EP98/00497</b>		(81) Designated States: AL, AU, BA, BB, BG, BR, CA, CN, CU, CZ, DE, EE, GE, GW, HU, ID, IL, IS, JP, KP, KR, LC, LK, LR, LT, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, SL, TR, TT, UA, US, UZ, VN, YU, ARIPO patent (GH, GM, KE, LS, MW, SD, SZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).	
(22) International Filing Date: 30 January 1998 (30.01.98)		<b>Published</b> <i>Without international search report and to be republished upon receipt of that report.</i>	
(30) Priority Data: 97101531.8 31 January 1997 (31.01.97) EP (34) Countries for which the regional or international application was filed: DE et al.			
(71) Applicant (for all designated States except US): BIOGNOSTIK GESELLSCHAFT FÜR BIOMOLEKULARE DIAGNOSTIK MBH [DE/DE]; Gerhard-Gerdes-Strasse 19, D-37079 Göttingen (DE).			
(72) Inventors; and (75) Inventors/Applicants (for US only): SCHLINGENSIEPEN, Karl-Hermann [DE/DE]; Pappelweg 3, D-37085 Göttingen (DE). BRYSCH, Wolfgang [DE/DE]; Calsowstrasse 56, D-37085 Göttingen (DE).			
(74) Agents: MEYERS, Hans-Wilhelm et al.; P.O. Box 10 22 41, D-50462 Cologne (DE).			
(54) Title: AN ANTISENSE OLIGONUCLEOTIDE PREPARATION METHOD			
(57) Abstract			
<p>A method for the preparation of an antisense oligonucleotide or derivative thereof comprising the steps of: selecting a target nucleic acid, if necessary elucidating its sequence; generating the antisense oligonucleotide with the proviso that: the oligonucleotide comprises at least 8 residues; the oligonucleotide comprises at maximum twelve elements, which are capable of forming three hydrogen bonds each to cytosine bases; the oligonucleotide does not contain four or more consecutive elements, capable of forming three hydrogen bonds each with four consecutive cytosine bases (CCCC) within the target molecule or alternatively four or more consecutive elements of GGGG; the oligonucleotide does also not contain 2 or more series of three consecutive elements, capable of forming three hydrogen bonds each with three consecutive cytosine bases (CCC) within the target molecule, or alternatively 2 or more series of three consecutive elements of GGG; and the ratio between residues forming two hydrogen bonds per residue (2H-bond-R) with the target molecule and those residues forming three hydrogen bonds per residue (3H-bond-R) with the target molecule, is ruled by the following specifications: 3H-bond-R/3H-bond-R + 2H-bond-R <math>\geq</math> 0.29; and synthesizing the oligonucleotide thus generated in a per se known manner.</p>			